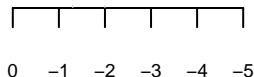
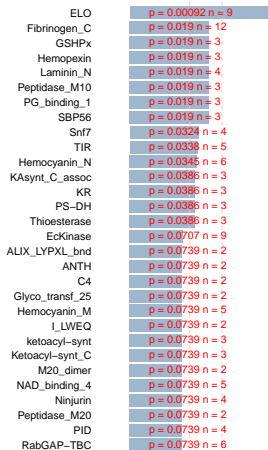


Fem.Oecy-Bulk_difpos domain



log(p)

n = 718 genes | p adjust BH TRUE